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RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/769,508

DATE: 02/07/2002
TIME: 13:43:36

Input Set : A:\Bebio111.app
Output Set: N:\CRF3\02072002\I769508.raw

ENTERED

```
 3 <110> APPLICANT: STUART, SUSAN G.
 4         MONAHAN, JOHN J.
 5         LANGTON, BEATRICE CLAUDIA
 6         HANCOCK, MIRIAM E.C.
 7         CHAO, LORRINE A.
 8         BLUFORD, PETER
10 <120> TITLE OF INVENTION: C-ERBB-2 EXTERNAL DOMAIN: GP75
12 <130> FILE REFERENCE: BEBIO-111-C1
14 <140> CURRENT APPLICATION NUMBER: 09/769,508
15 <141> CURRENT FILING DATE: 2001-01-26
17 <160> NUMBER OF SEQ ID NOS: 2
19 <170> SOFTWARE: PatentIn Ver. 2.1
21 <210> SEQ ID NO: 1
22 <211> LENGTH: 4543
23 <212> TYPE: DNA
24 <213> ORGANISM: Homo sapiens
26 <220> FEATURE:
27 <221> NAME/KEY: CDS
28 <222> LOCATION: (150)..(3914)
30 <400> SEQUENCE: 1
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33 cccggccccc acccctcgca gcacccccgcg cccccgccct cccagccggg tccagccgga 120
35 gccatggggc cggagccgca gtgagcacc atg gag ctg gcg gcc ttg tgc cgc   173
36                                   Met Glu Leu Ala Ala Leu Cys Arg
37                                    1               5
39 tgg ggg ctc ctc ctc gcc ctc ttg ccc ccc gga gcc gcg agc acc caa   221
40 Trp Gly Leu Leu Leu Ala Leu Leu Pro Pro Gly Ala Ala Ser Thr Gln
41      10                  15                  20
43 gtg tgc acc ggc aca gac atg aag ctg cgg ctc cct gcc agt ccc gag   269
44 Val Cys Thr Gly Thr Asp Met Lys Leu Arg Leu Pro Ala Ser Pro Glu
45  25                  30                  35                  40
47 acc cac ctg gac atg ctc cgc cac ctc tac cag ggc tgc cag gtg gtg   317
48 Thr His Leu Asp Met Leu Arg His Leu Tyr Gln Gly Cys Gln Val Val
49                  45                  50                  55
51 cag gga aac ctg gaa ctc acc tac ctg ccc acc aat gcc agc ctg tcc   365
52 Gln Gly Asn Leu Glu Leu Thr Tyr Leu Pro Thr Asn Ala Ser Leu Ser
53                  60                  65                  70
55 ttc ctg cag gat atc cag gag gtg cag ggc tac gtg ctc atc gct cac   413
56 Phe Leu Gln Asp Ile Gln Glu Val Gln Gly Tyr Val Leu Ile Ala His
57                  75                  80                  85
59 aac caa gtg agg cag gtc cca ctg cag agg ctg cgg att gtg cga ggc   461
60 Asn Gln Val Arg Gln Val Pro Leu Gln Arg Leu Arg Ile Val Arg Gly
61          90                  95                  100
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 63 acc cag ctc ttt gag gac aac tat gcc ctg gcc gtg cta gac aat gga      509
 64 Thr Gln Leu Phe Glu Asp Asn Tyr Ala Leu Ala Val Leu Asp Asn Gly
 65 105               110               115               120
 67 gac ccg ctg aac aat acc acc cct gtc aca ggg gcc tcc cca gga ggc      557
 68 Asp Pro Leu Asn Asn Thr Thr Pro Val Thr Gly Ala Ser Pro Gly Gly
 69               125               130               135
 71 ctg cgg gag ctg cag ctt cga agc ctc aca gag atc ttg aaa gga ggg      605
 72 Leu Arg Glu Leu Gln Leu Arg Ser Leu Thr Glu Ile Leu Lys Gly Gly
 73               140               145               150
 75 gtc ttg atc cag cgg aac ccc cag ctc tgc tac cag gac acg att ttg      653
 76 Val Leu Ile Gln Arg Asn Pro Gln Leu Cys Tyr Gln Asp Thr Ile Leu
 77         155               160               165
 79 tgg aag gac atc ttc cac aag aac aac cag ctg gct ctc aca ctg ata      701
 80 Trp Lys Asp Ile Phe His Lys Asn Asn Gln Leu Ala Leu Thr Leu Ile
 81     170               175               180
 83 gac acc aac cgc tct cgg gcc tgc cac ccc tgt tct ccg atg tgt aag      749
 84 Asp Thr Asn Arg Ser Arg Ala Cys His Pro Cys Ser Pro Met Cys Lys
 85 185               190               195               200
 87 ggc tcc cgc tgc tgg gga gag agt tct gag gat tgt cag agc ctg acg      797
 88 Gly Ser Arg Cys Trp Gly Glu Ser Ser Glu Asp Cys Gln Ser Leu Thr
 89               205               210               215
 91 cgc act gtc tgt gcc ggt ggc tgt gcc cgc tgc aag ggg cca ctg ccc      845
 92 Arg Thr Val Cys Ala Gly Gly Cys Ala Arg Cys Lys Gly Pro Leu Pro
 93               220               225               230
 95 act gac tgc tgc cat gag cag tgt gct gcc ggc tgc acg ggc ccc aag      893
 96 Thr Asp Cys Cys His Glu Gln Cys Ala Ala Gly Cys Thr Gly Pro Lys
 97         235               240               245
 99 cac tct gac tgc ctg gcc tgc ctc cac ttc aac cac agt ggc atc tgt      941
100 His Ser Asp Cys Leu Ala Cys Leu His Phe Asn His Ser Gly Ile Cys
101     250               255               260
103 gag ctg cac tgc cca gcc ctg gtc acc tac aac aca gac acg ttt gag      989
104 Glu Leu His Cys Pro Ala Leu Val Thr Tyr Asn Thr Asp Thr Phe Glu
105 265               270               275               280
107 tcc atg ccc aat ccc gag ggc cgg tat aca ttc ggc gcc agc tgt gtg     1037
108 Ser Met Pro Asn Pro Glu Gly Arg Tyr Thr Phe Gly Ala Ser Cys Val
109               285               290               295
111 act gcc tgt ccc tac aac tac ctt tct acg gac gtg gga tcc tgc acc     1085
112 Thr Ala Cys Pro Tyr Asn Tyr Leu Ser Thr Asp Val Gly Ser Cys Thr
113         300               305               310
115 ctc gtc tgc ccc ctg cac aac caa gag gtg aca gca gag gat gga aca     1133
116 Leu Val Cys Pro Leu His Asn Gln Glu Val Thr Ala Glu Asp Gly Thr
117         315               320               325
119 cag cgg tgt gag aag tgc agc aag ccc tgt gcc cga gtg tgc tat ggt     1181
120 Gln Arg Cys Glu Lys Cys Ser Lys Pro Cys Ala Arg Val Cys Tyr Gly
121     330               335               340
123 ctg ggc atg gag cac ttg cga gag gtg agg gca gtt acc agt gcc aat     1229
124 Leu Gly Met Glu His Leu Arg Glu Val Arg Ala Val Thr Ser Ala Asn
125 345               350               355               360
127 atc cag gag ttt gct ggc tgc aag aag atc ttt ggg agc ctg gca ttt     1277
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128 Ile Gln Glu Phe Ala Gly Cys Lys Lys Ile Phe Gly Ser Leu Ala Phe
129               365               370               375
131 ctg ccg gag agc ttt gat ggg gac cca gcc tcc aac act gcc ccg ctc   1325
132 Leu Pro Glu Ser Phe Asp Gly Asp Pro Ala Ser Asn Thr Ala Pro Leu
133               380               385               390
135 cag cca gag cag ctc caa gtg ttt gag act ctg gaa gag atc aca ggt   1373
136 Gln Pro Glu Gln Leu Gln Val Phe Glu Thr Leu Glu Glu Ile Thr Gly
137               395               400               405
139 tac cta tac atc tca gca tgg ccg gac agc ctg cct gac ctc agc gtc   1421
140 Tyr Leu Tyr Ile Ser Ala Trp Pro Asp Ser Leu Pro Asp Leu Ser Val
141         410               415               420
143 ttc cag aac ctg caa gta atc cgg gga cga att ctg cac aat ggc gcc   1469
144 Phe Gln Asn Leu Gln Val Ile Arg Gly Arg Ile Leu His Asn Gly Ala
145 425               430               435               440
147 tac tcg ctg acc ctg caa ggg ctg ggc atc agc tgg ctg ggg ctg cgc   1517
148 Tyr Ser Leu Thr Leu Gln Gly Leu Gly Ile Ser Trp Leu Gly Leu Arg
149               445               450               455
151 tca ctg agg gaa ctg ggc agt gga ctg gcc ctc atc cac cat aac acc   1565
152 Ser Leu Arg Glu Leu Gly Ser Gly Leu Ala Leu Ile His His Asn Thr
153               460               465               470
155 cac ctc tgc ttc gtg cac acg gtg ccc tgg gac cag ctc ttt cgg aac   1613
156 His Leu Cys Phe Val His Thr Val Pro Trp Asp Gln Leu Phe Arg Asn
157         475               480               485
159 ccg cac caa gct ctg ctc cac act gcc aac cgg cca gag gac gag tgt   1661
160 Pro His Gln Ala Leu Leu His Thr Ala Asn Arg Pro Glu Asp Glu Cys
161         490               495               500
163 gtg ggc gag ggc ctg gcc tgc cac cag ctg tgc gcc cgc agg gca ctg   1709
164 Val Gly Glu Gly Leu Ala Cys His Gln Leu Cys Ala Arg Arg Ala Leu
165 505               510               515               520
167 ctg ggg tca ggg ccc acc cag tgt gtc aac tgc agc cag ttc ctt cgg   1757
168 Leu Gly Ser Gly Pro Thr Gln Cys Val Asn Cys Ser Gln Phe Leu Arg
169               525               530               535
171 ggc cag gag tgc gtg gag gaa tgc cga gta ctg cag ggg ctc ccc agg   1805
172 Gly Gln Glu Cys Val Glu Glu Cys Arg Val Leu Gln Gly Leu Pro Arg
173              540               545               550
175 gag tat gtg aat gcc agg cac tgt ttg ccg tgc cac cct gag tgt cag   1853
176 Glu Tyr Val Asn Ala Arg His Cys Leu Pro Cys His Pro Glu Cys Gln
177         555               560               565
179 ccc cag aat ggc tca gtg acc tgt ttt gga ccg gag gct gac cag tgt   1901
180 Pro Gln Asn Gly Ser Val Thr Cys Phe Gly Pro Glu Ala Asp Gln Cys
181         570               575               580
183 gtg gcc tgt gcc cac tat aag gac cct ccc ttc tgc gtg gcc cgc tgc   1949
184 Val Ala Cys Ala His Tyr Lys Asp Pro Pro Phe Cys Val Ala Arg Cys
185 585               590               595               600
187 ccc agc ggt gtg aaa cct gac ctc tcc tac atg ccc atc tgg aag ttt   1997
188 Pro Ser Gly Val Lys Pro Asp Leu Ser Tyr Met Pro Ile Trp Lys Phe
189               605               610               615
191 cca gat gag gag ggc gca tgc cag cct tgc ccc atc aac tgc acc cac   2045
192 Pro Asp Glu Glu Gly Ala Cys Gln Pro Cys Pro Ile Asn Cys Thr His
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193             620               625                 630
195 tcc tgt gtg gac ctg gat gac aag ggc tgc ccc gcc gag cag aga gcc    2093
196 Ser Cys Val Asp Leu Asp Asp Lys Gly Cys Pro Ala Glu Gln Arg Ala
197         635                 640                 645
199 agc cct ctg acg tcc atc gtc tct gcg gtg gtt ggc att ctg ctg gtc    2141
200 Ser Pro Leu Thr Ser Ile Val Ser Ala Val Val Gly Ile Leu Leu Val
201   650                 655                 660
203 gtg gtc ttg ggg gtg gtc ttt ggg atc ctc atc aag cga cgg cag cag    2189
204 Val Val Leu Gly Val Val Phe Gly Ile Leu Ile Lys Arg Arg Gln Gln
205 665                 670                 675                 680
207 aag atc cgg aag tac acg atg cgg aga ctg ctg cag gaa acg gag ctg    2237
208 Lys Ile Arg Lys Tyr Thr Met Arg Arg Leu Leu Gln Glu Thr Glu Leu
209             685                 690                 695
211 gtg gag ccg ctg aca cct agc gga gcg atg ccc aac cag gcg cag atg    2285
212 Val Glu Pro Leu Thr Pro Ser Gly Ala Met Pro Asn Gln Ala Gln Met
213         700                 705                 710
215 cgg atc ctg aaa gag acg gag ctg agg aag gtg aag gtg ctt gga tct    2333
216 Arg Ile Leu Lys Glu Thr Glu Leu Arg Lys Val Lys Val Leu Gly Ser
217         715                 720                 725
219 ggc gct ttt ggc aca gtc tac aag ggc atc tgg atc cct gat ggg gag    2381
220 Gly Ala Phe Gly Thr Val Tyr Lys Gly Ile Trp Ile Pro Asp Gly Glu
221     730                 735                 740
223 aat gtg aaa att cca gtg gcc atc aaa gtg ttg agg gaa aac aca tcc    2429
224 Asn Val Lys Ile Pro Val Ala Ile Lys Val Leu Arg Glu Asn Thr Ser
225 745                 750                 755                 760
227 ccc aaa gcc aac aaa gaa atc tta gac gaa gca tac gtg atg gct ggt    2477
228 Pro Lys Ala Asn Lys Glu Ile Leu Asp Glu Ala Tyr Val Met Ala Gly
229             765                 770                 775
231 gtg ggc tcc cca tat gtc tcc cgc ctt ctg ggc atc tgc ctg aca tcc    2525
232 Val Gly Ser Pro Tyr Val Ser Arg Leu Leu Gly Ile Cys Leu Thr Ser
233         780                 785                 790
235 acg gtg cag ctg gtg aca cag ctt atg ccc tat ggc tgc ctc tta gac    2573
236 Thr Val Gln Leu Val Thr Gln Leu Met Pro Tyr Gly Cys Leu Leu Asp
237     795                 800                 805
239 cat gtc cgg gaa aac cgc gga cgc ctg ggc tcc cag gac ctg ctg aac    2621
240 His Val Arg Glu Asn Arg Gly Arg Leu Gly Ser Gln Asp Leu Leu Asn
241     810                 815                 820
243 tgg tgt atg cag att gcc aag ggg atg agc tac ctg gag gat gtg cgg    2669
244 Trp Cys Met Gln Ile Ala Lys Gly Met Ser Tyr Leu Glu Asp Val Arg
245 825                 830                 835                 840
247 ctc gta cac agg gac ttg gcc gct cgg aac gtg ctg gtc aag agt ccc    2717
248 Leu Val His Arg Asp Leu Ala Ala Arg Asn Val Leu Val Lys Ser Pro
249             845                 850                 855
251 aac cat gtc aaa att aca gac ttc ggg ctg gct cgg ctg ctg gac att    2765
252 Asn His Val Lys Ile Thr Asp Phe Gly Leu Ala Arg Leu Leu Asp Ile
253         860                 865                 870
255 gac gag aca gag tac cat gca gat ggg ggc aag gtg ccc atc aag tgg    2813
256 Asp Glu Thr Glu Tyr His Ala Asp Gly Gly Lys Val Pro Ile Lys Trp
257         875                 880                 885
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259 atg gcg ctg gag tcc att ctc cgc cgg cgg ttc acc cac cag agt gat      2861
260 Met Ala Leu Glu Ser Ile Leu Arg Arg Arg Phe Thr His Gln Ser Asp
261     890                 895                 900
263 gtg tgg agt tat ggt gtg act gtg tgg gag ctg atg act ttt ggg gcc      2909
264 Val Trp Ser Tyr Gly Val Thr Val Trp Glu Leu Met Thr Phe Gly Ala
265 905                 910                 915                 920
267 aaa cct tac gat ggg atc cca gcc cgg gag atc cct gac ctg ctg gaa      2957
268 Lys Pro Tyr Asp Gly Ile Pro Ala Arg Glu Ile Pro Asp Leu Leu Glu
269                 925                 930.                935
271 aag ggg gag cgg ctg ccc cag ccc ccc atc tgt acc att gat gtc tac      3005
272 Lys Gly Glu Arg Leu Pro Gln Pro Pro Ile Cys Thr Ile Asp Val Tyr
273             940                 945                 950
275 atg atc atg gtc aaa tgt tgg atg att gac tct gaa tgt cgg cca aga      3053
276 Met Ile Met Val Lys Cys Trp Met Ile Asp Ser Glu Cys Arg Pro Arg
277         955                 960                 965
279 ttc cgg gag ttg gtg tct gaa ttc tcc cgc atg gcc agg gac ccc cag      3101
280 Phe Arg Glu Leu Val Ser Glu Phe Ser Arg Met Ala Arg Asp Pro Gln
281     970                 975                 980
283 cgc ttt gtg gtc atc cag aat gag gac ttg ggc cca gcc agt ccc ttg      3149
284 Arg Phe Val Val Ile Gln Asn Glu Asp Leu Gly Pro Ala Ser Pro Leu
285 985                 990                 995             1000
287 gac agc acc ttc tac cgc tca ctg ctg gag gac gat gac atg ggg gac      3197
288 Asp Ser Thr Phe Tyr Arg Ser Leu Leu Glu Asp Asp Asp Met Gly Asp
289                 1005                1010                1015
291 ctg gtg gat gct gag gag tat ctg gta ccc cag cag ggc ttc ttc tgt      3245
292 Leu Val Asp Ala Glu Glu Tyr Leu Val Pro Gln Gln Gly Phe Phe Cys
293             1020                1025                1030
295 cca gac cct gcc ccg ggc gct ggg ggc atg gtc cac cac agg cac cgc      3293
296 Pro Asp Pro Ala Pro Gly Ala Gly Gly Met Val His His Arg His Arg
297         1035                1040                1045
299 agc tca tct acc agg agt ggc ggt ggg gac ctg aca cta ggg ctg gag      3341
300 Ser Ser Ser Thr Arg Ser Gly Gly Gly Asp Leu Thr Leu Gly Leu Glu
301     1050                1055                1060
303 ccc tct gaa gag gag gcc ccc agg tct cca ctg gca ccc tcc gaa ggg      3389
304 Pro Ser Glu Glu Glu Ala Pro Arg Ser Pro Leu Ala Pro Ser Glu Gly
305 1065                1070                1075.               1080
307 gct ggc tcc gat gta ttt gat ggt gac ctg gga atg ggg gca gcc aag      3437
308 Ala Gly Ser Asp Val Phe Asp Gly Asp Leu Gly Met Gly Ala Ala Lys
309             1085                1090                1095
311 ggg ctg caa agc ctc ccc aca cat gac ccc agc cct cta cag cgg tac      3485
312 Gly Leu Gln Ser Leu Pro Thr His Asp Pro Ser Pro Leu Gln Arg Tyr
313             1100                1105                1110
315 agt gag gac ccc aca gta ccc ctg ccc tct gag act gat ggc tac gtt      3533
316 Ser Glu Asp Pro Thr Val Pro Leu Pro Ser Glu Thr Asp Gly Tyr Val
317         1115                1120                1125
319 gcc ccc ctg acc tgt agc ccc cag cct gaa tat gtg aac cag cca gat      3581
320 Ala Pro Leu Thr Cys Ser Pro Gln Pro Glu Tyr Val Asn Gln Pro Asp
321     1130                1135                1140
323 gtt cgg ccc cag ccc cct tcg ccc cga gag ggc cct ctg cct gct gcc      3629
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